



SEQUENCE LISTING

<110> Brennan, Thomas J.
Leviten, Michael W.

<120> TRANSGENIC MICE CONTAINING CERBERUS GENE
DISRUPTIONS

<130> R-67

<140> US 09/887,552

<141> 2001-06-21

<150> US 60/213,670

<151> 2000-06-21

<150> US 60/266,046

<151> 2001-02-01

<150> US 60/282,668

<151> 2001-04-09

<160> 4

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 1752

<212> DNA

<213> Mus musculus

<220>

<221> misc_feature

<222> 1235, 1313

<223> n = A,T,C or G

<400> 1

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tctgggaaag gcagacctat gtgtggatgg ctgccagagt cagggctctt tatcctttcc 180
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gaggcagaga gggaagatgc tgtccaggct tggaagattc tggaagaaac ctgagaccga 360
attttacccc ccaagggatg tggaaagcga tcatgtctca tcggggatgc aggccgtgac 420
tcagccagca gatgggagga aagtggagag atcacctcta caggaggaag ccaagagggt 480
ctggcatcgg ttcattgttca gaaagggccc ggcgttccag ggagtcattc tgcccatcaa 540
aagccacgaa gtacactggg agacctgcag gactgtgccc ttcaaccaga ccattgcccc 600
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aattacctca acagaaagca aaacctcaac agaataagtg aggggttattc aatctggaaa 960
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tcctgagtgc tgggattaca gacatgctcc ataanacata gctcccagaa ggatttttaa 1260
aagagatttt gcatgtttca aagttgcctt tgagactcag aaatatcttg atntattgaa 1320
tggccttgcc acagatgtgg gaggcagctt gcttggtggc ccaagtattt tttttttgtt 1380
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cgttcagaat tctccacatg aagtttttac tgttggttat ctggcgttga agaaggaata 1440
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tcccgggaata tgaggaaaat acatgaacag tattagagtc acttgaggaa gttactagga 1560
aacgccataa gtctccaagt acattgtgag tcattttgaa ggacaatcgt gtatatagac 1620
gaaatcttct actcgtatgc ttttgaatct tctagcaagt taggtttcta tgtttgggct 1680
tcttcctatt gtctaagagt atgtgtgaca aattcaacct gacaaatacc tcaatggcaa 1740
attctgaccc tg 1752

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<210> 2

<211> 272

<212> PRT

<213> Mus musculus

<400> 2

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Met His Leu Leu Leu Val Gln Leu Leu Val Leu Leu Pro Leu Gly Lys
 1          5          10          15
Ala Asp Leu Cys Val Asp Gly Cys Gln Ser Gln Gly Ser Leu Ser Phe
 20          25          30
Pro Leu Leu Glu Arg Gly Arg Arg Asp Leu His Val Ala Asn His Glu
 35          40          45
Glu Ala Glu Asp Lys Pro Asp Leu Phe Val Ala Val Pro His Leu Met
 50          55          60
Gly Thr Ser Leu Ala Gly Glu Gly Gln Arg Gln Arg Gly Lys Met Leu
 65          70          75          80
Ser Arg Leu Gly Arg Phe Trp Lys Lys Pro Glu Thr Glu Phe Tyr Pro
 85          90          95
Pro Arg Asp Val Glu Ser Asp His Val Ser Ser Gly Met Gln Ala Val
 100         105         110
Thr Gln Pro Ala Asp Gly Arg Lys Val Glu Arg Ser Pro Leu Gln Glu
 115         120         125
Glu Ala Lys Arg Phe Trp His Arg Phe Met Phe Arg Lys Gly Ala Pro
 130         135         140
Phe Gln Gly Val Ile Leu Pro Ile Lys Ser His Glu Val His Trp Glu
 145         150         155         160
Thr Cys Arg Thr Val Pro Phe Asn Gln Thr Ile Ala His Glu Asp Cys
 165         170         175
Gln Lys Val Val Val Gln Asn Asn Leu Cys Phe Gly Lys Cys Ser Ser
 180         185         190
Ile Arg Phe Pro Gly Glu Gly Ala Asp Ala His Ser Phe Cys Ser His
 195         200         205
Cys Ser Pro Thr Lys Phe Thr Thr Val His Leu Met Leu Asn Cys Thr
 210         215         220
Ser Pro Thr Pro Val Val Lys Met Val Met Gln Val Glu Glu Cys Gln
 225         230         235         240
Cys Met Val Lys Thr Glu Arg Gly Glu Glu Arg Leu Leu Leu Ala Gly
 245         250         255
Ser Gln Gly Ser Phe Ile Pro Gly Leu Pro Ala Ser Lys Thr Asn Pro
 260         265         270

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<210> 3

<211> 200

<212> DNA

<213> Artificial Sequence

<220>

<223> Targeting vector

<400> 3

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cagctgcttg ttctcttgcc tctggggaag gcagacctat gtgtggatgg ctgccagagt 120

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```

cagggctctt tatcctttcc tctcccagaa aggggtcgca gagatctcca cgtggccaac 180
cacgaggagg cagaagacaa                                200

```

```

<210> 4
<211> 200
<212> DNA
<213> Artificial Sequence

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<220>
<223> Targeting vector

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<400> 4
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aggacaaacg gcaaaataga aagagtctgg cgagagctcg ggccttgtct agttccagat 180
tcagtccttt gggatttcat                                200

```